

Turner

PT-12

1644

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/181,311

DATE: 02/22/2000
TIME: 15:31:48

Input Set: RECEIVED

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This Raw Listing contains the General Information
Section and up to first 5 pages.

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1 <110> APPLICANT: Lee, Arthur M.E.
2 Jain, Mukesh
3 Watanabe, Masafumi
4 <120> TITLE OF INVENTION: IN VITRO DIFFERENTIATION OF VASCULAR SMOOTH MUSCLE
5 CELLS AND REAGENTS RELATED THERETO
6 <130> FILE REFERENCE: APV-382.01
7 <140> CURRENT APPLICATION NUMBER: US/09/181,311
8 <141> CURRENT FILING DATE: 1998-10-28
9 <150> EARLIER APPLICATION NUMBER: 60/063,363
10 <151> EARLIER FILING DATE: 1997-10-28
11 <150> EARLIER APPLICATION NUMBER: 60/080,420
12 <151> EARLIER FILING DATE: 1998-04-02
13 <150> EARLIER APPLICATION NUMBER: 60/096,685
14 <151> EARLIER FILING DATE: 1998-08-14
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16 <170> SOFTWARE: PatentIn Ver. 2.0
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18 <211> LENGTH: 153
19 <212> TYPE: DNA
20 <213> ORGANISM: Murine
21 <220> FEATURE:
22 <223> OTHER INFORMATION: Clone WO11
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51	50	55	60	
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54	Ala	Glu	Val	Pro
55	85	90	95	
56	Lys	Asp	Lys	Gly
57	100	105	110	
58	Arg	Pro	Pro	Lys
59	115	120	125	
60	Pro	Lys	Glu	Lys
61	130	135	140	
62	Pro	Lys	Ala	Thr
63	145	150	155	160
64	Pro	Ser	Gly	Lys
65	165	170	175	
66	Trp	Pro	Leu	Pro
67	180	185	190	
68	Glu	Gly	Gly	Ala
69	195	200	205	
70	Thr	His	Val	Glu
71	210	215	220	
72	Glu	Gln	Pro	Thr
73	225	230	235	240
74	Phe	Glu	Tyr	Ile
75	245	250	255	
76	Arg	Arg	Arg	Pro
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78	Pro	Ala	Pro	Ala
79	275	280	285	
80	Leu	Pro	Pro	Leu
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82	Tyr	Asp	Asp	Met
83	305	310	315	320
84	Glu	Arg	Gln	Thr
85	325	330	335	
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87	340	345	350	
88	Gly	Lys	Asp	His
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90	Trp	Thr	Pro	Thr
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92	His	Arg	Ile	Asn
93	385	390	395	400
94	Gly	Ala	Gln	Arg

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100	Thr Gln Gly Arg Asp Ser Ser Ile His Asp Asp Phe Val Thr Thr Phe						
101		450		455		460	
102	Phe Phe Ser Asn Asp Ser Gln Thr Trp Val Met Tyr Thr Asn Gly Tyr						
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106	Ser Glu Leu Pro Glu Pro Val Val Ala Arg Phe Ile Arg Ile Tyr Pro						
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110	Ser Val Ala Pro Val Tyr Ser Tyr Tyr Ala Gln Asn Glu Val Val Asp						
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112	Asp Leu Asp Phe Arg His His Ser Tyr Lys Asp Met Arg Gln Leu Met						
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115		565		570		575	
116	Gly Lys Ser Ser Arg Gly Leu Lys Ile Tyr Ala Met Glu Ile Ser Asp						
117		580		585		590	
118	Asn Pro Gly Glu His Glu Leu Gly Glu Pro Glu Phe Arg Tyr Thr Ala						
119		595		600		605	
120	Gly Ile His Gly Asn Glu Val Leu Gly Arg Glu Leu Ile Leu Met Gln						
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122	Tyr Leu Cys Arg Glu Tyr Arg Asp Gly Asn Pro Arg Val Arg Ser Leu						
123		625		630		635	640
124	Val Gln Asp Thr Arg Ile His Leu Val Pro Ser Leu Asn Pro Asp Gly						
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126	Tyr Glu Val Ala Ala Gln Met Gly Ser Glu Phe Gly Asn Trp Ala Leu						
127		660		665		670	
128	Gly Leu Trp Thr Glu Glu Gly Phe Asp Ile Phe Glu Asp Phe Pro Asp						
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130	Leu Asn Ser Val Leu Trp Gly Ala Glu Glu Phe Val Pro Tyr Arg Val						
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132	Pro Asn Asn Asn Leu Pro Ile Pro Glu Arg Tyr Leu Ser Pro Asp Ala						
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134	Thr Val Ser Thr Glu Val Arg Ala Ile Ile Ala Trp Met Glu Lys Asn						
135		725		730		735	
136	Pro Phe Val Leu Gly Ala Asn Leu Asn Gly Gly Glu Arg Leu Val Ser						
137		740		745		750	
138	Tyr Pro Tyr Asp Met Ala Arg Thr Pro Thr Gln Glu Gln Leu Leu Ala						
139		755		760		765	
140	Ala Ala Met Ala Ala Ala Glu Gly Glu Asp Glu Val Ser Glu Ala Gln						
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142	Glu Thr Pro Asp His Ala Ile Phe Arg Trp Leu Ala Ile Ser Phe Ala						
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144	Ser Ala His Leu Thr Leu Thr Glu Pro Tyr Arg Gly Gly Cys Gln Ala						

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149          835          840          845
150  Leu Glu Leu Ser Phe Tyr Ile Asp Lys Phe Pro His Glu Ser Glu Leu
151          850          855          860
152  Pro Arg Glu Trp Glu Asn Asn Lys Glu Ala Leu Leu Thr Phe Met Glu
153          865          870          875          880
154  Gln Val His Arg Gly Ile Lys Gly Val Val Thr Asp Glu Gln Gly Ile
155          885          890          895
156  Pro Ile Ala Asn Ala Thr Ile Ser Val Ser Gly Ile Asn His Gly Val
157          900          905          910
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159          915          920          925
160  Tyr Arg Val Thr Ala Glu Gly Tyr Thr Pro Ser Ala Lys Thr Cys Asn
161          930          935          940
162  Val Asp Tyr Asp Ile Gly Ala Thr Gln Cys Asn Phe Ile Leu Ala Arg
163          945          950          955          960
164  Ser Asn Trp Lys Arg Ile Arg Glu Ile Met Ala Met Asn Gly Asn Arg
165          965          970          975
166  Pro Ile Pro His Ile Asp Pro Ser Arg Pro Met Thr Pro Gln Gln Arg
167          980          985          990
168  Arg Leu Gln Gln Arg Arg Leu Gln His Arg Leu Arg Leu Arg Ala Gln
169          995          1000          1005
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177          1060          1065          1070
178  Glu Phe Gly Thr Lys Val Glu Pro Glu Phe Glu Thr Gln Leu Phe Glu
179          1075          1080          1085
180  Thr Gln Leu Glu Pro Glu Phe Glu Glu Glu Glu Glu Glu Lys Glu
181          1090          1095          1100
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209	Lys Glu Lys Pro Pro Lys Ala Thr Lys Lys Pro Lys Glu Lys Pro Pro	130	135	140
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213	Ala Gly Lys Lys Phe Ser Thr Val Ala Pro Leu Glu Thr Leu Asp Arg	165	170	175
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218				
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220				
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223	Glu Tyr Ile Arg Arg Gln Lys Gln Pro Arg Pro Thr Pro Ser Arg Arg	245	250	255
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226				
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228				
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231	Pro Pro Pro Gln Lys Pro Asp Val Gly Gln Glu Val Asp Glu Glu Lys	305	310	315
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237	Arg Lys Gly Glu Glu Leu Glu Glu Glu Trp Ala Pro Val Glu Lys Ile	355	360	365
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243	Asn Met Gln Ala Gly Ala Asn Glu Asp Asp Tyr Tyr Asp Gly Ala Trp	405	410	415
244				

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VERIFICATION SUMMARY
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Line ? Error/Warning

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